## SEQUENGE LISTING

						~					-10				
	<b>9</b> 110	>	Tor	nen	Corp	pora	itio	n/							
500	<del>~120</del> :	>	Met	hod	foi	c Me	asy	reme	ent	of	hepa	atit	is	C vi	irus
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	Phe M	et Gly	Thr	Asn	Pro	Lys	Pro	Gln	Arg	Lys	Thr	Lys	Arg	Asn	Thr
	_		20	/				25					30		
	Asn A	rg Arg 35	Pro	Gl	Asp	Val		Phe	Pro	Gly	Gly		Gln	Ile	Val
•	Glv G	ly Val	Tvr	T	T.e.u	Pro	40 Ara	Δτα	G1 v	Pro	Ara	45	Gl v	Va 1	Δ×α
		50	-1-	$\mathcal{T}^{-}$		55	9	9	OL,		60	200	GLY	141	nig
÷	Ala T	hr Arg	Lys	Thr	Ser		Arg	Ser	Gln	Pro		Gly	Gly	Arg	Arg
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	Pro I	le Pro	Lys	Asp	Arg	Arg	Ser	Thr	Gly	Lys	Ser	Trp	Gly	Lys	Pro
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	M T	T	100		_		_	105	_	_	_		110		_
	Trp L	eu Leu 115	ser	Pro	Arg	СТĀ	120	Arg	Pro	Ser	Trp	_	Pro	Thr	Asp
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                         135
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Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Phe Arg Val Gly Ala Phe
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Thr Arg Lys Thr Ser Lys Arg/Ser Gln Pro Arg Gly Gly Arg Arg Pro
Ile Pro Lys Asp Arg Ser Thr Gly Lys Ser Trp Gly Lys Pro Gly
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                                         75
                                                              80
Tyr Pro Trp Pro Leu Ty
                         Sly Asn Glu Gly Leu Gly Trp Ala Gly Trp
Leu Leu Ser Pro Arg/Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
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                                105
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Arg His Arg Ser Arg Asn Val Gly Lys Val Ile Asp Thr Leu Thr Cys
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Leu Val Leu	Asp Pro Ser Val Ala Ser Thr Leu Gly Phe Gly Ala Tyr
	20 25 30
ctg age aag	gcc cat ggt gtg aac ccg aac atc cgc acg ggc atc cgt 144

Ä

Let	ı Se	r Lys	s Ala	a His	Gly	/ Val	Asr	Pro	Ası	ıle	Arq	y Thr	Gly	, Ile	Arg	
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Va.	l Gl	y Tì	r P	ro Ly	ys Se	r Ar	g Ar	g Pr	o Gl	u Gly	y Ar	a Ala	a Tr	o Al	a C	21 n	
225	5				23					23		-				240	
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Pro	G1;	y Th	r Il	.e Il	e Le	u Se	r Gly	/ Ar	g Pro	o Ala	a Val	l Val	l Pro	) As	n A	ıra	, 00
				24					250					25		_ 9	
gaa	gt	g ct	g ta	t ca	a ga	a tti	cto	gaç	ggc	tct	aga:	a gcc	g get			tt	816
Glu	Va.	l Le	u Ty	r Gl	n Gl	u Phe	Let	ı Glı	ı Ala	a Ser	Arc	, Ala	a Ala	Lei	u I	le	020
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Glu	Gli	ı Gl	y Gl	n Ar	g Ile	a Ala	Glu	Met	Lev	ı Lys	Ser	Lys	: Ile	Glr	a G	lv	
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tta	ctg	ca	g ca	a gc	c tco	aag	cag	gcc	caa	gac	ata	aaa	ato	gad		a+	912
Leu	Leu	Gl	n Gl	n Ala	a Ser	: Lys	Gln	Ala	Gln	Asp	Ile	Lvs	Ile	Ast	. G	lv	312
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acc	ctg	at	t at	t cc	g aaa	gat	cgt	cgc	age	acc	ggt	aaa	agc	taa	rac	at.	960
Thr	Leu	Ile	∍ Ile	e Pro	o Lys	Asp	Arg	Arg	Ser	Thr	Gly	Lys	Ser	Tro	. GI	lv	300
305					310					315	-	-			32		
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Lys	Pro	Gl	Phe	e Leu	ı Ile	Asp	Ser	Leu	His	Ile	Asn	Gln	Ara	Ala	. Va	11	1000
				325					330					335			
gtt	gca	ccg	gac	aag	gag	gtc	ctt	tat	gag	gct	ttt	gat	σaσ			ď	1056
Val	Ala	Pro	Asp	Lys	Glu	Val	Leu	Tyr	Glu	Ala	Phe	Asp	Glu	Met	Gl	11	1030
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ctc	gcc	atg	ggc	acc	aac	ccg	aaa	ccg	gag	cgt	aaa	agc	aag	cat	aa	c	1104
Leu	Ala	Met	Gly	Thr	Asn	Pro	Lys	Pro	Glu	Arg	Lys	Ser	Lys	Ara	As	n	-
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acc	aac	cgt	aaa	ccg	cag	gat	att	aaa	ttc	ccg	ggt	agt	ggt	cag	at	ar a	1152
Thr.	Asn	Arg	Lys	Pro	Gln	Asp	Ile	Lys	Phe	Pro	Gly	Ser	Gly	Gln	Va.	1	
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Le	ı Va	l Lei	Asp 20	Pro	Ser	'Val	L Ala	Ser 25	Thr	Leu	Gly	Phe	e Gl <sub>3</sub>		a Tyr
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Thr	7 Va]	Thr	Thr	: Gly	Ala	Pro	Val	Thr	Tyr	Ser	Thr	Туг	Gly	y Lys	Tyr
Leu 65	ı Ala	Asp	Gly	Gly	C <b>y</b> s 70	Ala	Gly	Gly	Ala	<b>Tyr</b> 75	Asp	Val	Ile	Gly	Ser 80
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Cys	His	Ser 115	Lys	Glu	Lys	Cys	Asp 120	Glu	Leu	Ala	Ser	Ala			Gly
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Thr	Gly	Phe	Thr	Gly 165	Asp	Phe	Asp	Ser	Val 170	Val	Asp	Cys	Asn	Thr 175	
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Ile	Glu	Gln 195	Gly	Met	Gln	Leu	Ser 200	Glu	Gln	Phe	Lys	Gln 205		Ser	Leu
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Val 225	Gly	Thr	Pro		Ser . 230	Arg	Arg	Pro		Gly 235	Arg	Ala	Trp	Ala	Gln 240

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Glu	Val	Leu	Tyr	Gln	Glu	Phe	Leu	Glu	Ala	Ser	Arg	Ala	Ala	Leu	Ile
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